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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/826,025**

**DATE: 12/18/2001**  
**TIME: 16:38:14**

**Input Set : N:\CrF3\RULE60\09826025.raw**  
**Output Set: N:\CRF3\12182001\I826025.raw**

**SEQUENCE LISTING**

- 3 (1) GENERAL INFORMATION:  
 5     (i) APPLICANT: Chang, Lung-Ji  
 7     (ii) TITLE OF INVENTION: Combination Immunogene Therapy  
 9     (iii) NUMBER OF SEQUENCES: 25  
 11    (iv) CORRESPONDENCE ADDRESS:  
 12      (A) ADDRESSEE: Medlen & Carroll, LLP  
 13      (B) STREET: 220 Montgomery Street, Suite 2200  
 14      (C) CITY: San Francisco  
 15      (D) STATE: California  
 16      (E) COUNTRY: United States of America  
 17      (F) ZIP: 94104  
 19    (v) COMPUTER READABLE FORM:  
 20      (A) MEDIUM TYPE: Floppy disk  
 21      (B) COMPUTER: IBM PC compatible  
 22      (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
 23      (D) SOFTWARE: PatentIn Release #1.0, Version #1.30  
 25    (vi) CURRENT APPLICATION DATA:  
 C--> 26      (A) APPLICATION NUMBER: US/09/826,025  
 C--> 27      (B) FILING DATE: 04-Apr-2001  
 28      (C) CLASSIFICATION:  
 30    (vii) PRIOR APPLICATION DATA:  
 31      (A) APPLICATION NUMBER: 08/838,702  
 32      (B) FILING DATE:  
 35    (viii) ATTORNEY/AGENT INFORMATION:  
 36      (A) NAME: Ingolia, Diane E.  
 37      (B) REGISTRATION NUMBER: 40,027  
 38      (C) REFERENCE/DOCKET NUMBER: CHANG-02687  
 40    (ix) TELECOMMUNICATION INFORMATION:  
 41      (A) TELEPHONE: (415) 705-8410  
 42      (B) TELEFAX: (415) 397-8338  
 45 (2) INFORMATION FOR SEQ ID NO: 1:  
 47    (i) SEQUENCE CHARACTERISTICS:  
 48      (A) LENGTH: 6145 base pairs  
 49      (B) TYPE: nucleic acid  
 50      (C) STRANDEDNESS: double  
 51      (D) TOPOLOGY: linear  
 53    (ii) MOLECULE TYPE: DNA (genomic)  
 58    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
 60 GAATTCTAC CAGATCACCG AAAACTGTCC TCCAAATGTG TCCCCCTCAC ACTCCCAAAT       60  
 62 TCGCGGGCTT CTGCCTCTTA GACCACTCTA CCCTATTCCC CACACTCACC GGAGCCAAAG       120  
 64 CGCGGGCCCT TCCGTTCTT TGCTTTGAA AGACCCCACC CGTAGGTGGC AAGCTAGCTT       180  
 66 AAGTAACGCC ACTTTGCAAG GCATGGAAAA ATACATAACT GAGAATAGAA AAGTTCAGAT       240  
 68 CAAGGTCAGG AACAAAGAAA CAGCTGAATA CCAAACAGGA TATCTGTGGT AAGCGGTTCC       300  
 70 TGCCCCGGCT CAGGGCCAAG AACAGATGAG ACAGCTGAGT GATGGGCCAA ACAGGATATC       360  
 72 TGTGGTAAGC AGTTCTGCC CCGGCTCGGG GCCAAGAAC AATGGTCCCC AGATGCCGTC       420  
 74 CAGCCCTCAG CAGTTCTAG TGAATCATCA GATGTTCCA GGGTGCCCCA AGGACCTGAA       480
- ENTERED**

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76	AATGACCCCTG	TACCTTATTG	GAACTAACCA	ATCAGTCGC	TTCTCGCTTC	TGTCGCGCG	540
78	CTTCCGCTCT	CCGAGCTCAA	AAAAAGAGCC	CACAACCCCT	CACTCGGCCG	GCCAGTCTTC	600
80	CGATAGACTG	CGTCGCCCCG	GTACCCGTAT	TCCCAATAAA	GCCTCTTGCT	GTTTGCATCC	660
82	GAATCGTGGT	CTCGCTGTT	CTTGGGAGGG	TCTCCTCTGA	GTGATTGACT	ACCCACGACG	720
84	GGGGTCTTTC	ATTGGGGGC	TCGTCCGGGA	TTTGGAGACC	CCTGCCAGG	GACCACCGAC	780
86	CCACCAACGG	GAGGTAAGCT	GGCCAGCAAC	TTATCTGTGT	CTGTCCGATT	GTCTAGTGTG	840
88	TATGTTTGAT	GTTATGCGCC	TGCGTCTGTA	CTAGTTAGCT	AACTAGCTCT	GTATCTGGCG	900
90	GACCCGTGGT	GGAACTGACG	AGTTCTGAAC	ACCCGGCCGC	AACCTGGGA	GACGTCCCAG	960
92	GGACTTTGGG	GGCCGTTTT	GTGGCCCGAC	CTGAGGAAGG	GAGTCGATGT	GGAAATCCGAC	1020
94	CCCGTCAGGA	TATGTGGTTC	TGGTAGGAGA	CGAGAACCTA	AAACAGTTC	CGCCTCCGTC	1080
96	TGAATTTTG	CTTTCGGTT	CGAACCGAAG	CCGCCGCGTCT	TGTCGCTGC	AGCGCTGCAG	1140
98	CATCGTTCTG	TGTTGCTCT	GTCTGACTGT	GTTCCTGTAT	TGTCGAA	ATTAGGGCCA	1200
100	GACTGTTACC	ACTCCCTAA	GTTCGACCTT	AGGTCACTGG	AAAGATGTCG	AGCGGATCGC	1260
102	TCACAACCAAG	TCGGTAGATG	TCAAGAAGAG	ACGTTGGGTT	ACCTTCTGCT	CTGCAGAATG	1320
104	GCCAACCTT	AACGTGGAT	GGCCGCGAGA	CGGCACCTT	AACCGAGACC	TCATCACCCA	1380
106	GGTTAACATC	AAGGTCTTT	CACCTGGCCC	GCATGGACAC	CCAGACCAGG	TCCCCTACAT	1440
108	CGTGACCTGG	GAAGCCTTGG	CTTTGACCC	CCCTCCCTGG	GTCAAGCCT	TTGTACACCC	1500
110	TAAGCCTCCG	CCTCCCTTTC	CTCCATCCGC	CCCGTCTCTC	CCCCTTGAAAC	CTCCTCGTTC	1560
112	GACCCCGCCT	CGATCCTCCC	TTTATCCAGC	CCTCACTCCT	TCTCTAGGCG	CCGGAATTCC	1620
114	GATCTGATCA	AGAGACAGGA	TGAGGATCGT	TTCGCATGAT	TGAACAAGAT	GGATTGCACG	1680
116	CAGGTTCTCC	GGCCGCTTGG	GTGGAGAGGC	TATTCGGCTA	TGACTGGCA	CAACAGACAA	1740
118	TCGGCTGCTC	TGATGCCGCC	GTGTTCCGGC	TGTCAGCGCA	GGGGCGCCCG	GTTCTTTTG	1800
120	TCAAGACCGA	CCTGTCCGGT	GCCCTGAATG	AACTGCAGGA	CGAGGCAGCG	CGGCTATCGT	1860
122	GGCTGGCAC	GACGGCGTT	CCTTGCAG	CTGTGCTCGA	CGTTGTCACT	GAAGCGGGAA	1920
124	GGGACTGGCT	GCTATTGGGC	GAAGTGCCTGG	GGCAGGATCT	CCTGTATCT	CACCTTGCTC	1980
126	CTGCCGAGAA	AGTATCCATC	ATGGCTGATG	CAATGCCCG	GCTGCATACG	CTTGATCCGG	2040
128	CTACCTGCC	ATTGACCCAC	CAAGCGAAAC	ATCGCATCGA	GCGAGCACGT	ACTCGGATGG	2100
130	AAGCCGGTCT	TGTCGATCAG	GATGATCTGG	ACGAAGAGCA	TCAGGGCTC	GCGCCAGCCG	2160
132	AACTGTTCGC	CAGGCTCAAG	GCGCGCATGC	CCGACGGCGA	GGATCTCGTC	GTGACCCATG	2220
134	GCGATGCCCTG	CTTGCGGAAT	ATCATGGTGG	AAAATGGCCG	CTTTTCTGGA	TTCATCGACT	2280
136	GTGGCCGGCT	GGGTGTGGCG	GACCGCTATC	AGGACATAGC	GTGCTTAC	CGTGATATTG	2340
138	CTGAAGAGCT	TGGCGCGAA	TGGGCTGACC	GCTTCCCTCGT	GCTTACGGT	ATCGCCGCTC	2400
140	CCGATTGCGA	GCGCATCGCC	TTCTATCGCC	TTCTGACGA	GTTCTTCTGA	CGGGGACTCT	2460
142	GGGGTTCGAA	ATGACCGACC	AAGCGACGCC	CAACCTGCCA	TCACGAGATT	TCGATTCCAC	2520
144	CGCCGCTTC	TATGAAAGGT	TGGGCTTCGG	AATCGTTTC	CGGGACGCCG	GCTGGATGAT	2580
146	CCTCCAGCGC	GGGGATCTCA	TGCTGGAGTT	CTTCGCCAC	CCCGGGCTCG	ATCCCCTCGC	2640
148	GAGTTGGTTC	AGCTGCTGCC	TGAGGCTGGA	CGACTCGCG	GAGTTCTACC	GGCAGTGCAA	2700
150	ATCCGTCGGC	ATCCAGGAAA	CCAGCAGCGG	CTATCCGCG	ATCCATGCC	CCGAACGTGCA	2760
152	GGAGTGGGGA	GGCACGATGG	CCGCTTTGGT	CGACCCGGAC	GGGACGCTCC	TGCGCCTGAT	2820
154	ACAGAACGAA	TTGCTGCA	GCATCTCATG	AGTGTGCTT	CCCCTTTCC	GCCTGAGGTC	2880
156	ACTGCGTGG	TGGAGCGCTG	GGCGCTGCTG	CGCGACGGCG	AGCTGCTCAC	CACCCACTCG	2940
158	AGGGCGTGC	GCGCTGCAGA	GGCCGAGTGC	AGAACTGCTC	CAAAGGGACC	TCAAGGCTTT	3000
160	CCGAGGGACA	CTAGGCTGAC	TCCATCGAGC	CAGTGTAGAG	ATAAGCTTAT	CGATTAGTCC	3060
162	AATTGTTAA	AGACAGGATA	TCAGTGGTCC	AGGCTCTAGT	TTTGACTCAA	CAATATCACC	3120
164	AGCTGAAGCC	TATAGAGTAC	GAGCCATAGA	TAAAATAAAA	GATTTTATT	AGTCTCCAGA	3180
166	AAAAGGGGGG	AATGAAAGAC	CCCACCTGTA	GGTTGGCAA	GCTAGCTTAA	GTAACGCCAT	3240
168	TTTGCAGGGC	ATGGAAAAAT	ACATAACTGA	GAATAGAGAA	GTTCAGATCA	AGGTCAGGAA	3300
170	CAGATGGAAC	AGCTGAATAT	GGGCCAAACA	GGATATCTGT	GGTAAGCAGT	TCCTGCCCG	3360
172	GCTCAGGGCC	AAGAACAGAT	GGAACAGCTG	AATATGGGCC	AAACAGGATA	TCTGTGGTAA	3420

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174	GCAGTTCTG	CCCCGGCTCA	GGGCCAAGAA	CAGATGGTCC	CCAGATGCGG	TCCAGCCCTC	3480
176	AGCAGTTCT	AGAGAACCAT	CAGATGTTTC	CAGGGTGCCC	CAAGGACCTG	AAATGACCCCT	3540
178	GTGCCTTATT	TGAACTAACC	AATCAGTTCG	CTTCTCGCTT	CTGTTCGCGC	GCTTCTGCTC	3600
180	CCCGAGCTCA	ATAAAAGAGC	CCACAACCCC	TCACTCGGGG	CGCCAGTCCT	CCGATTGACT	3660
182	GAGTCGCCCC	GGTACCCGTG	TATCCAATAA	ACCCCTCTGC	AGTTGCATCC	GACTTGTGGT	3720
184	CTCGCTGTT	CTTGGGAGGG	TCTCCTCTGA	GTGATTGACT	ACCCGTCA	GGGGGTCTTT	3780
186	CATTTGGGGG	CTCGTCCGGG	ATCGGGAGAC	CCCTGCCAG	GGACCACCGA	CCCACCAACCG	3840
188	GGAGGTAAGC	TGGCTGCCTC	GCGCGTTCG	GTGATGACGG	TGAAAACCTC	TGACACATGC	3900
190	AGCTCCCAGA	GACGGTCACA	GCTTGTCTGT	AAGCGGATGC	CGGGAGCAGA	CAAGCCCCTC	3960
192	AGGGCGCGTC	AGCGGGTGT	GGCGGGTGT	GGGGCGCAGC	CATGACCCAG	TCACGTAGCG	4020
194	ATAGCGGAGT	GTATACTGGC	TTAACTATGC	GGCATCAGAG	CAGATTGTAC	TGAGAGTGCA	4080
196	CCATATGCCG	TGTGAAATAC	CGCACAGATG	CGTAAGGAGA	AAATACCGCA	TCAGGCGCTC	4140
198	TTCCGCTTCC	TCGCTCACTG	ACTCGCTGCG	CTCGGTGTT	CGGCTGCCGC	GAGCGGTATC	4200
200	AGCTCACTCA	AAGCGGTAA	TACGGTTATC	CACAGAATCA	GGGGATAACG	CAGGAAAGAA	4260
202	CATGTGAGCA	AAAGGCCAGC	AAAAGGCCAG	GAACCGTAA	AAGGCCGCGT	TGCTGGCGTT	4320
204	TTTCCATAGG	CTCCGCC	CTGACGAGCA	TCACAAAAAT	CGACGCTCAA	GTCAGAGGTG	4380
206	GCGAAACCCG	ACAGGACTAT	AAAGATACCA	GGCGTTCCC	CCTGGAAGCT	CCCTCGTGC	4440
208	CTCTCCTGTT	CCGACCCCTGC	CGCTTACCGG	ATACCTGTCC	GCCTTCCTCC	CTTCGGGAAG	4500
210	CGTGGCGCTT	TCTCATAGCT	CACGCTGAG	GTATCTCAGT	TCGGTGTAGG	TCGTTGCTC	4560
212	CAAGCTGGGC	TGTGTGCACG	AACCCCCCGT	TCAGCCCGAC	CGCTGCGCCT	TATCCGGTAA	4620
214	CTATCGTCTT	GAGTCCAACC	CGGTAAGACA	CGACTTATCG	CCACTGGCAG	CAGCCACTGG	4680
216	TAACAGGATT	AGCAGAGCGA	GGTATGTAGG	CGGGTCTACA	GAGTTCTGA	AGTGGTGGCC	4740
218	TAACATCGGC	TACACTAGAA	GGACAGTATT	TGGTATCTGC	GCTCTGCTGA	AGCCAGTTAC	4800
220	CTTCGGAAAA	AGAGTTGGTA	GCTCTTGATC	CGGCAAACAA	ACCACCGCTG	GTAGCGGTGG	4860
222	TTTTTTGTT	TGCAAGCAGC	AGATTACCGC	CAGAAAAAAA	GGATCTCAAG	AAGATCCTT	4920
224	GATCTTTCT	ACGGGGTCTG	ACGCTCAGTG	GAACGAAAAC	TCACGTTAAG	GGATTGGT	4980
226	CATGAGATTA	TCAAAAAGGA	TCTTCACCTA	GATCCTTTA	AATTAAAAAT	GAAGTTTTAA	5040
228	ATCAATCTAA	AGTATATATG	AGTAAACTTG	GTCTGACAGT	TACCAATGCT	TAATCAGTGA	5100
230	GGCACCTATC	TCAGCGATCT	GTCTATTTCG	TTCATCCATA	GTTGCCGTAC	TCCCCGTCGT	5160
232	GTAGATAACT	ACGATACGGG	AGGGCTTAC	ATCTGGCCCC	AGTGTGCAA	TGATACCGCG	5220
234	AGACCCACGC	TCACCGGCTC	CAGATTATC	AGCAATAAAC	CAGCCAGCG	GAAGGGCCGA	5280
236	GCGCAGAAGT	GGTCCTGCAA	CTTTATCCGC	CTCCATCCAG	TCTATTAAATT	TTGCCGGGA	5340
238	AGCTAGAGTA	AGTAGTTCGC	CAGTTAATAG	TTTGGCIAAC	GTTGTTGCCA	TTGCTGCAGG	5400
240	CATCGTGGTG	TCACGCTCGT	CGTTTGGTAT	GGCTTCATTC	AGCTCCGGTT	CCCAACGATC	5460
242	AAGGCGAGTT	ACATGATCCC	CCATGTTGTG	CAAAAAAGCG	GTTAGCTCCT	TCGGTCCCTCC	5520
244	GATCGTTGTC	AGAAGTAAGT	TGGCCGCAGT	GTTATCACTC	ATGGTTATGG	CAGCACTGCA	5580
246	TAATTCTCTT	ACTGTCA	CATCCGTAAG	ATGCTTTCT	GTGACTGGT	AGTACTCAAC	5640
248	CAAGTCATTC	TGAGAAATAGT	GTATGCGGG	ACCGAGTTGC	TCTTGGCCGG	CGTCAACACG	5700
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252	GGGGCGAAAA	CTCTCAAGGA	TCTTACCGCT	GTTGAGATCC	AGTTGATGT	AACCCACTCG	5820
254	TGCACCAAC	TGATCTTCAG	CATCTTTAC	TTTCACCGC	GTTTCTGGT	GAGCAAAAC	5880
256	AGGAAGGCAA	AATGCCGAA	AAAAGGGAAT	AAGGGCGACA	CGGAAATGTT	GAATACTCAT	5940
258	ACTCTCCTT	TTTCAATATT	ATTGAAGCAT	TTATCAGGGT	TATTGTCTCA	TGAGCGGATA	6000
260	CATATTGAA	TGTATTAGA	AAAATAAAC	AATAGGGTT	CCGCGCACAT	TTCCCCGAAA	6060
262	AGTGCCACCT	GACGTCTAAG	AAACCATTAT	TATCATGACA	TTAACCTATA	AAAATAGGCG	6120
264	TATCACGAGG	CCCTTCGTC	TTC				6145

266 (2) INFORMATION FOR SEQ ID NO: 2:

268 (i) SEQUENCE CHARACTERISTICS:  
269 (A) LENGTH: 67 base pairs

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/826,025

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Input Set : N:\Crf3\RULE60\09826025.raw  
 Output Set: N:\CRF3\12182001\I826025.raw

270	(B) TYPE: nucleic acid	
271	(C) STRANDEDNESS: single	
272	(D) TOPOLOGY: linear	
274	(ii) MOLECULE TYPE: other nucleic acid	
275	(A) DESCRIPTION: /desc = "DNA"	
280	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
282	GATCTAAGCT TCGGGCCGCA GATCTCGAGC CATGGATCCT AGGCCTGATC ACGCGTCGAC	60
284	TCGCGAT	67
286	(2) INFORMATION FOR SEQ ID NO: 3:	
288	(i) SEQUENCE CHARACTERISTICS:	
289	(A) LENGTH: 65 base pairs	
290	(B) TYPE: nucleic acid	
291	(C) STRANDEDNESS: single	
292	(D) TOPOLOGY: linear	
294	(ii) MOLECULE TYPE: other nucleic acid	
295	(A) DESCRIPTION: /desc = "DNA"	
300	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
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304	GCTTA	65
306	(2) INFORMATION FOR SEQ ID NO: 4:	
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309	(A) LENGTH: 33 base pairs	
310	(B) TYPE: nucleic acid	
311	(C) STRANDEDNESS: single	
312	(D) TOPOLOGY: linear	
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315	(A) DESCRIPTION: /desc = "DNA"	
320	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
322	AAGCTTGATC ACCACCATGA TTGAACAAGA TGG	33
324	(2) INFORMATION FOR SEQ ID NO: 5:	
326	(i) SEQUENCE CHARACTERISTICS:	
327	(A) LENGTH: 34 base pairs	
328	(B) TYPE: nucleic acid	
329	(C) STRANDEDNESS: single	
330	(D) TOPOLOGY: linear	
332	(ii) MOLECULE TYPE: other nucleic acid	
333	(A) DESCRIPTION: /desc = "DNA"	
338	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
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342	(2) INFORMATION FOR SEQ ID NO: 6:	
344	(i) SEQUENCE CHARACTERISTICS:	
345	(A) LENGTH: 35 base pairs	
346	(B) TYPE: nucleic acid	
347	(C) STRANDEDNESS: single	
348	(D) TOPOLOGY: linear	
350	(ii) MOLECULE TYPE: other nucleic acid	
351	(A) DESCRIPTION: /desc = "DNA"	
356	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
358	CCCGGGAAGC TTCCACCATG TGGCTGCAGA GCCTG	35

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/826,025 DATE: 12/18/2001  
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360 (2) INFORMATION FOR SEQ ID NO: 7:  
 362 (i) SEQUENCE CHARACTERISTICS:  
 363 (A) LENGTH: 29 base pairs  
 364 (B) TYPE: nucleic acid  
 365 (C) STRANDEDNESS: single  
 366 (D) TOPOLOGY: linear  
 368 (ii) MOLECULE TYPE: other nucleic acid  
 369 (A) DESCRIPTION: /desc = "DNA"  
 374 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:  
 376 AATGGATCCT ATCACTCCTG GACTGGCTC 29  
 378 (2) INFORMATION FOR SEQ ID NO: 8:  
 380 (i) SEQUENCE CHARACTERISTICS:  
 381 (A) LENGTH: 435 base pairs  
 382 (B) TYPE: nucleic acid  
 383 (C) STRANDEDNESS: double  
 384 (D) TOPOLOGY: linear  
 386 (ii) MOLECULE TYPE: other nucleic acid  
 387 (A) DESCRIPTION: /desc = "DNA"  
 392 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:  
 394 ATGTGGCTGC AGAGCCTGCT GCTCTGGGC ACTGTGGCCT GCAGCATCTC TGCACCCGCC 60  
 396 CGCTCGCCA GCCCCAGCAC GCAGCCCTGG GAGCATGTGA ATGCCATCCA GGAGGCCGG 120  
 398 CGTCTCCTGA ACCTGAGTAG AGACACTGCT GCTGAGATGA ATGAAACAGT AGAAGTCATC 180  
 400 TCAGAAATGT TTGACCTCCA GGAGCCGACC TGCCCTACAGA CCCGCCTGGA GCTGTACAAG 240  
 402 CAGGGCCTGC GGGCAGCCT CACCAAGCTC AAGGGCCCT TGACCATGAT GGCCAGCCAC 300  
 404 TACAAGCAGC ACTGCCCTCC AACCCCGGAA ACTTCCGTG CAACCCAGAT TATCACCTT 360  
 406 GAAAGTTTCA AAGAGAACCT GAAGGACTTT CTGCTTGTCA TCCCCTTGGA CTGCTGGGAG 420  
 408 CCAGTCCAGG AGTGA 435  
 410 (2) INFORMATION FOR SEQ ID NO: 9:  
 412 (i) SEQUENCE CHARACTERISTICS:  
 413 (A) LENGTH: 30 base pairs  
 414 (B) TYPE: nucleic acid  
 415 (C) STRANDEDNESS: single  
 416 (D) TOPOLOGY: linear  
 418 (ii) MOLECULE TYPE: other nucleic acid  
 419 (A) DESCRIPTION: /desc = "DNA"  
 424 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
 426 TGTGGATCCA CCATGGGACT GAGTAACATT 30  
 428 (2) INFORMATION FOR SEQ ID NO: 10:  
 430 (i) SEQUENCE CHARACTERISTICS:  
 431 (A) LENGTH: 35 base pairs  
 432 (B) TYPE: nucleic acid  
 433 (C) STRANDEDNESS: single  
 434 (D) TOPOLOGY: linear  
 436 (ii) MOLECULE TYPE: other nucleic acid  
 437 (A) DESCRIPTION: /desc = "DNA"  
 442 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:  
 444 TTTGGATCCT TAAAAACATG TATCACTTT GTCGC 35  
 446 (2) INFORMATION FOR SEQ ID NO: 11:  
 448 (i) SEQUENCE CHARACTERISTICS:

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/826,025

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TIME: 16:38:15

Input Set : N:\Crf3\RULE60\09826025.raw  
Output Set: N:\CRF3\12182001\I826025.raw

L:26 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:27 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]



✓ 03c0

I hereby certify that this correspondence is being deposited with the United States Postal Service as first class mail in an envelope addressed to:  
Assistant Commissioner for Patents  
Washington, D.C. 20231 on June 27, 2001.

David Saliwanchik  
David R. Saliwanchik, Patent Attorney

REQUEST TO USE CRF FROM PRIOR APPLICATION AND STATEMENT  
Examining Group  
Patent Application  
Docket No. CNG-100D1  
Serial No. 09/826,025

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Art Unit : 1632  
Applicant : Lung-Ji Chang  
Serial No. : 09/826,025  
Filed : April 4, 2001  
For : Combination Immunogene Therapy

Box SEQUENCE

Assistant Commissioner for Patents  
Washington, D.C. 20231

REQUEST TO USE COMPUTER READABLE FORMAT FROM  
PRIOR APPLICATION AND STATEMENT UNDER 37 CFR §1.821

Sir:

It is respectfully requested that the computer readable format of patent application Serial No. 08/838,702 entitled "Combination Immunogene Therapy," filed April 9, 1997 by the applicant, Lung-Ji Chang, also be used as the computer readable format for the above-identified patent application. I hereby certify that the sequence listing in the subject application and the sequence listing in application Serial No. 08/838,702 contain identical sequence information and that the paper and computer readable copies contain the same information.

A Notice to Comply with Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures was received from the Patent Office, and a copy of that Notice is attached herewith.

Respectfully submitted,



David R. Saliwanchik  
Patent Attorney  
Registration No. 31,794  
Phone No.: 352-375-8100  
Fax No.: 352-372-5800  
Address: 2421 N.W. 41st Street, Suite A-1  
Gainesville, FL 32606-6669

DRS/sl

Attachment: copy of Notice to Comply with Requirements and/or Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures